

SEQUENCE LISTING

	1.	SEQ ID NO:1:C101	CCGAGTGAACCTGCGGAAA
	2.	SEQ ID NO:2:C102	TACTACAACCAGAGCGAGGA
5	3.	SEQ ID NO:3:C103	CACAGACTGACCGAGTGAG
	4.	SEQ ID NO:4:C104	AGTCCAAGAGGGGAGCCG
	5.	SEQ ID NO:5:C105	CCACTCCATGAGGTATTTCT
	6.	SEQ ID NO:6:C106	TCTTCTCCAGAAGGCACCAT
	7.	SEQ ID NO:7:C107	CAGGTCAGTGTGATCTCCA
10	8.	SEQ ID NO:8:C108	CCTCCAGGTAGGCTCTCCA
	9.	SEQ ID NO:9:C109	CAGCCCCTCGTGCTGCAT
	10.	SEQ ID NO:10:C110	CGCGCGCTGCAGCGTCTT
	11.	SEQ ID NO:11:C111	CCTCCAGGTAGGCTCTCAG
	12.	SEQ ID NO:12:C112	CTCAGGGTGAGGGGCTCT
15	13.	SEQ ID NO:13:C113	TGAGCCGCCGTGTCCGCA
	14.	SEQ ID NO:14:C114	GGTCGCAGCCATACATCCA
	15.	SEQ ID NO:15:C115	CCGCGGGTATGACCAGTC
	16.	SEQ ID NO:16:C116	GCGTCTCCTTCCCGTTCTT
	17.	SEQ ID NO:17:C117	AGCGTCTCCTTCCCATTCTT
20	18.	SEQ ID NO:18:C118	TCCGCGGGTATG ACCAGTA
	19.	SEQ ID NO:19:C119	GCCCCAGGTCGCAGCCAA
	20.	SEQ ID NO:20:C120	ACAAGCGCCAGGCACAGG
	21.	SEQ ID NO:21:C121	GAGCCACTCCACGCACTC
	22.	SEQ ID NO:22:C122	CCCTCCAGGTAGGCTCTCT
25	23.	SEQ ID NO:23:C123	TCGTAGGCTAACTGGTCATG
	24.	SEQ ID NO:24:C124	CCGCCGTGTCCGCGGCA
	25.	SEQ ID NO:25:C125	TACAACCAGAGCGAGGCCA
	26.	SEQ ID NO:26:C126	ACAACCAGAGCGAGGCCG
	27.	SEQ ID NO:27:C127	ACGACACGCAGTTCGTGCA
30	28.	SEQ ID NO:28:C128	GCGCAGGTTCCGCAGGC
	29.	SEQ ID NO:29:C129	GAGCCACTCCACGCACCG
	30.	SEQ ID NO:30:C130	GAGCCACTCCACGCACGT
	31.	SEQ ID NO:31:C131	CCTCCAGGTAGGCTCTCTG

	32. SEQ ID NO:32:C132	CCGCGGAGGAAGCGCCA
	33. SEQ ID NO:33:C133	CCACTCCATGAGGTATTTCTT
	34. SEQ ID NO:34:C134	CCGGAGTATTGGGACCTGC
	35. SEQ ID NO:35:C135	CCCCAGGTCGCAAGCCAG
5	36. SEQ ID NO:36:C136	CGCACGGGCGCCTCCA
	37. SEQ ID NO:37:C137	GCGCCGTGGATAGAGCAA
	38. SEQ ID NO:38:C138	GCCGCGAGTCCGAGGAC
	39. SEQ ID NO:39:C139	ACCGGAACACACAGATCTG
	40. SEQ ID NO:40:C140	ACCGGGAGACACAGATCTC
10	41. SEQ ID NO:41:C141	GGAGTATTGGGACCGGAAC
	42. SEQ ID NO:42:C142	AACATGAAGGCCTCCGCG
	43. SEQ ID NO:43:C143	GACCGGAACACACAGATCTT
	44. SEQ ID NO:44:C144	TACCGAGAGAACCTGCGC
	45. SEQ ID NO:45:C145	AGCAGGAGGGGCGCGAA
15	46. SEQ ID NO:46:C146	GGGGAGCCCCGCTTCATT
	47. SEQ ID NO:47:C147	CAGATCTACAAGGCCAGG
	48. SEQ ID NO:48:C148	CCATGAGGTATTTCTACACCG
	49. SEQ ID NO:49:C149	GACCGGAACACACAGATCTA
	50. SEQ ID NO:50:C150	CCGAGAGAGCCTGCGGGAA
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	53. SEQ ID NO:55:C155	CCTCCAGGTAGGCTCTGTC
	54. SEQ ID NO:56:C156	GAGGAGGCGCCCGTCG
	55. SEQ ID NO:57:C157	CTTGCCGTCGTAGGCGG
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	57. SEQ ID NO:59:C159	CGTTCAGGGCGATGTAATCT
	58. SEQ ID NO:60:C160	CGTGCCCTCCAGGTAGGT
	59. SEQ ID NO:61:C161	GAGCCACTCCACGCACTC
	60. SEQ ID NO:62:C162	CCAGGTATCTGCGGAGCG
30	61. SEQ ID NO:63:C163	CCGCGCGCTCCAGCGTG
	62. SEQ ID NO:64:C164	TACCAGCGCGCTCCAGCT
	63. SEQ ID NO:65:C165	GCCATACATCCTCTGGATGA
	64. SEQ ID NO:66:C166	CGTCGCAGCCATACATCAC
	65. SEQ ID NO:67:C167	CTCTCAGCTGCTCCGCCT

	66. SEQ ID NO:68:C168	GTCGTAGGCGGACTGGTC
	67. SEQ ID NO:69:C169	TCGTAGGCGTCCTGGTGG
	68. SEQ ID NO:70:C170	CTCCAACCTGCGCTGGGA
	69. SEQ ID NO:71:C171	GTGTGTTCCGGTCCCAATAT
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	72. SEQ ID NO:74:C174	CCATACATCGTCTGCCAA
	73. SEQ ID NO:75:C175	GCGCAGGTTCCGCAGGC
	74. SEQ ID NO:76:C176	GAGCCACTCCACGCACAG
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	76. SEQ ID NO:78:C178	GAGACACAGAAGTACAAGCG
	77. SEQ ID NO:79:C179	GCCGCGGTCCAGGAGCT
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	79. SEQ ID NO:81:C181	CGCGAGTCCGAGGATGGC
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	81. SEQ ID NO:83:C183	CCACTCCCATGAGGTATTTCC
	82. SEQ ID NO:84:C184	GCGGCGGTCCAGGAGCG
	83. SEQ ID NO:85:C185	CCTCCAGGTAGGCTCTCAA
	84. SEQ ID NO:86:C186	GCAGGTTCCGCAGGCTCT
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	93. SEQ ID NO:95:C195	CTCGGTCAGTCTGTGCCTT
	94. SEQ ID NO:96:C196	TCTCGGTAAGTCTGTGCCTT
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	96. SEQ ID NO:98:C198	CGTCGTAGGCGTACTGGTC
	97. SEQ ID NO:99:C199	CGACGCCGCGAGCCAGAA
	98. SEQ ID NO:100:CI100	GAGCCCGTCCACGCACTC
	99. SEQ ID NO:101:CI101	TCACAGACTGACCGAGCGAA

	100. SEQ ID NO:102:CI102	ACGGAATGTGAAGGCCCCAG
	101. SEQ ID NO:103:CI103	AGCGACGCCGCGAGCCA
	102. SEQ ID NO:104:CI104	GGCCGGAGTATTGGGACGA
	103. SEQ ID NO:105:CI105	GATAGAGCAGGAGAGGCCT
5	104. SEQ ID NO:106:CI106	TCACAGACTGACCGAGAGAG
	105. SEQ ID NO:107:CI107	CCCGGCCCGGCAGTGGA
	106. SEQ ID NO:108:CI108	GTGGATAGAGCAGGAGGGT
	107. SEQ ID NO:109:CI109	AGTTAATCCTTGCCGTCGTAA
	108. SEQ ID NO:110:CI110	CACTCCACGCACGTGCCA
10	109. SEQ ID NO:111:CI111	AGCGCAGGTCCTCGTTCAA
	110. SEQ ID NO:112:CI112	CCGTCGTAGGCGTGCTGT
	111. SEQ ID NO:113:CI113	CCAAGAGCGCAGGTCCTCT
	112. SEQ ID NO:114:CI114	ACACAGATCTACAAGACCAAC
	113. SEQ ID NO:115:CI115	GGACCCGGGAGACACAGAAC
15	114. SEQ ID NO:116:CI116	CCCCAGGTCGCAGCCAC
	115. SEQ ID NO:117:CI117	TCTCAGCTGCTCCGCCGT
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	117. SEQ ID NO:119:CI119	CCGAGTGAACCTGCGGAAA
	118. SEQ ID NO:120:CI120	TACTACAACCAGAGCGAGGA
20	119. SEQ ID NO:121:CI121	CACGACTGACCGAGTGAG
	120. SEQ ID NO:122:CI122	AGTCCAAGAGGGGAGCCG
	121. SEQ ID NO:123:CI123	CCACTCCATGAGGTATTTCTC
	122. SEQ ID NO:124:CI124	CCTCCAGGTAGGCTCTCCA
	123. SEQ ID NO:125:CI125	CAGCCCCTCGTGCTGCAT
25	124. SEQ ID NO:126:CI126	CGCGCGCTGCAGCGTCTT
	125. SEQ ID NO:127:CI127	CCTCCAGGTAGGCTTCAG
	126. SEQ ID NO:128:CI128	GGTCGCAGCCAAACATCCA
	127. SEQ ID NO:129:CI129	AGCGTCTCCTTCCCATTCTT
	128. SEQ ID NO:130:CI130	TACCGAGAGAACCTGCGCA
30	129. SEQ ID NO:131:CI131	CCTTGCCGTCGTAGGCGA
	130. SEQ ID NO:132:CI132	GTCGTAGGCGTCCTGGTC
	131. SEQ ID NO:133:CI133	CCACGTCGCAGCCATACATT
	132. SEQ ID NO:134:CI134	GCCGCGAGTTCGAGAGG
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	134. SEQ ID NO:136:CI136	GCCTTCACATTCCGTGTGTT
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	141. SEQ ID NO:143:CI143	CTTCACATTCCGTGTCTCCT
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	145. SEQ ID NO:147:CI147	GACGACACGCTGTTTCGTGA
	146. SEQ ID NO:148:CI150	ACGTCGCAGCCGTACATG
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	149. SEQ ID NO:151:C2F25A	CACTCCATGAGGTATTTCTGA
	150. SEQ ID NO:152:C2F25C	CACTCCATGAGGTATTTCTC
	151. SEQ ID NO:153:C2F32C	TGAGGTATTTCTACACCGCC
	152. SEQ ID NO:154:C3R195G	CCTCCAGGTAGGCTCTGTC
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	154. SEQ ID NO:156:C34076A	CCTTGCCGTCGTAGGCGT
	155. SEQ ID NO:157:C34076C	CCTTGCCGTCGTAGGCGG
	156. SEQ ID NO:158:C3R076T	CCTTGCCGTCGTAGGCGA
	157. SEQ ID NO:159:C3R075TA	CCTTGCCGTCGTAGGCTA
25	158. SEQ ID NO:160:C2F216A	TACAAGCGCCAGGCACAGA
	159. SEQ ID NO:161:CI53	ATGATGTTGACCTTTCCAGGG
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	161. SEQ ID NO:163:CI148	TGCCAAGTGGAGCACCCAA
	162. SEQ ID NO:164:CI149	GCATCTTGCTCTGTGCAGA
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	164. SEQ ID NO:166:CICptA2	GATGGAGCCGCGGTGGATAGAGCAAGGAGGG
	165. SEQ ID NO:167:CICptB1	CAGTTCGTGAGGTTCGACAGCGACGCC
	166. SEQ ID NO:168:CICptB2	CTGCGCGGCTACTACAACCAGAGCGAGGCC
	167. SEQ ID NO:169:DQ01	TCC[CT]CGCAGAGGATTTTCGTG

	168. SEQ ID NO:170:DQ02	GGAGCGCGTGCGGGG
	169. SEQ ID NO:171:DQ03	ACGGAGCGCGTGCGTCT
	170. SEQ ID NO:172:DQ04	GGACGGAGCGCGTGCGTTA
	171. SEQ ID NO:173:DQ05	GTACTCCTCTCGGTTATAGATGTG
5	172. SEQ ID NO:174:DQ06	GATCTCTTCTCGGTTATAGATGC
	173. SEQ ID NO:175:DQ07	GTCGCTGTCGAAGCGCA
	174. SEQ ID NO:176:DQ08	TGACGCCGCTGGGGCC
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10	177. SEQ ID NO:179:DQ11	GCTGTTCCAGTACTCGGCAA
	178. SEQ ID NO:180:DQ12	CAACTCCGCCCGGGTCCT
	179. SEQ ID NO:181:DQ13	GAAGGACATCCTGGAGAGGAA
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	185. SEQ ID NO:187:DR04	CACGTTTCTTGCAGCAGGA
	186. SEQ ID NO:188:DR05	CACGTTTCTTGGAGCTGCG
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	243. SEQ ID NO:245:DR62	GGCTGTTCCAGTACTCAGCG
	244. SEQ ID NO:246:DR63	GCTGTTCCAGTACTCGGCGA
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	246. SEQ ID NO:248:DR65	CCGCCTCTGCTCCAGGAG
	247. SEQ ID NO:249:DR66	CCGCGCCTGCTCCAGGAT
	248. SEQ ID NO:250:DR67	ACCGCGGCGCGCCTGTCT
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	251. SEQ ID NO:253:DR70	CACCTCGGCCCGCCTCC
	252. SEQ ID NO:254:DR71	GTCCACCGCGGCGCGCGT
	253. SEQ ID NO:255:DR72	TGTCCACCGCGGCCCGCT
	254. SEQ ID NO:256:DR73	TCCACCGCGGCCCGCGC
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	257. SEQ ID NO:259:DR76	TAGGTGTCCACCGCGGCG
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	266. SEQ ID NO:268:DR85	CAGCACGTTTCTTGGAGCTGT
	267. SEQ ID NO:269:DR86	TTCTTGTGGCAGCTTAAGTTTGA
	268. SEQ ID NO:270:DPA-E(PC)	GATCCCCCTGAGGTGACCGTG
	269. SEQ ID NO:271:DPA-F(PC)	CTGGGCCCCGGGGGTCATGGCC

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272. SEQ ID NO:274:DQCPT3	CACGTCGCTGTCTGAAGCGTGCGTACTCCTC
273. SEQ ID NO:275:DQCPT4	CACGTCGCTGTCTGAAGCGCGCGTACTCCTC
5 274. SEQ ID NO:276:DQCPT5	CACGTCGCTGTCTGAAGCGCACGTCTCCTC
275. SEQ ID NO:277:DRCPT1	DRCP TGGCGTGGGCGAGGCAGGGTAACCTTCTTA

10

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